



SEQUENCE LISTING

<110> O'DOWD, BRIAN F.
GEORGE, SUSAN R.

<120> METHOD OF IDENTIFYING TRANSMEMBRANE PROTEIN-INTERACTING COMPOUNDS

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<141> 2003-04-11

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Thr Leu Leu

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Glu Leu Leu

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Phe Asp Asn Pro
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Ile Thr

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Ile Thr

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Trp	Pro	Gly	Ile	Pro
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Thr	Val	Leu	Ala	Leu	Leu	Ser	His	Lys	Lys	Phe	Lys	Arg	Xaa	Lys	Ile
1				5					10					15	

Trp	Pro	Gly	Ile	Pro
			20	

<210> 111
 <211> 40

<212> DNA
 <213> Artificial Sequence

 <220>
 <223> primer

 <400> 111
 gctcttcggg ctcgagcagc gatgcgaccc tccgggacgg 40

<210> 112
 <211> 39
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> primer

 <400> 112
 ctatcctccg tggtaaccgct gctccaataa attcactgc 39

<210> 113
 <211> 37
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> primer

 <400> 113
 cacatcgttc ggaagaagtt taagcggagg ctgctgc 37

<210> 114
 <211> 40
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> primer

 <400> 114
 cctgcagcag cctccgctta aacttcttcc gaacgatgtg 40

<210> 115
 <211> 19
 <212> PRT
 <213> Artificial Sequence

 <220>
 <223> synthesized

 <400> 115

Arg Arg Arg His Ile Val Arg Lys Arg Thr Leu Arg Arg Leu Leu Gln
 1 5 10 15

Glu Arg Glu

<210> 116
<211> 19
<212> PRT
<213> Artificial Sequence

<220>
<223> synthesized

<400> 116

Arg Arg Arg His Ile Val Arg Lys Lys Phe Lys Arg Arg Leu Leu Gln
1 5 10 15

Glu Arg Glu

<210> 117
<211> 49
<212> DNA
<213> Artificial Sequence

<220>
<223> primer

<400> 117
gaggactctg aacaccgaat tcgccgccat ggacgggact gggctggtg

49

<210> 118
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
<223> primer

<400> 118
gtgtggcagg attcatctgg gtaccgcggt tgggtgctga ccggt

45

<210> 119
<211> 41
<212> DNA
<213> Artificial Sequence

<220>
<223> primer

<400> 119
cctctgagga cctgaaaaag aagagaaagg ctggcatcgc c

41

<210> 120
<211> 41
<212> DNA
<213> Artificial Sequence

<220>
<223> primer

<400> 120
ggcgatgccg gcctttctct tctttttcag gtcctcagag g

41

<210> 121
<211> 33
<212> PRT
<213> Artificial Sequence

<220>
<223> synthesized

<400> 121

Asn Pro Ile Ile Tyr Ala Phe Asn Ala Asp Phe Arg Lys Ala Phe Ser
1 5 10 15

Thr Leu Leu Ser Ser Glu Asp Leu Lys Lys Glu Glu Ala Ala Gly Ile
20 25 30

Ala

<210> 122
<211> 33
<212> PRT
<213> Artificial Sequence

<220>
<223> synthesized

<400> 122

Asn Pro Ile Ile Tyr Ala Phe Asn Ala Lys Lys Phe Lys Arg Phe Ser
1 5 10 15

Thr Leu Leu Ser Ser Glu Asp Leu Lys Lys Lys Arg Lys Ala Gly Ile
20 25 30

Ala

<210> 123
<211> 45
<212> DNA
<213> Artificial Sequence

<220>
<223> primer

<400> 123
cctagtcgcg agcaggccga attcgccacc atggacagca gcacc

45

<210> 124
 <211> 44
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> primer

 <400> 124
 gatggtgtga gaccggtacc gcgggcaatg gagcagtttc tgcc 44

 <210> 125
 <211> 45
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> primer

 <400> 125
 cctagtccgc agcaggccga attcgccacc atggacagca gcacc 45

 <210> 126
 <211> 45
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> primer

 <400> 126
 ggatggtgtg agaccggtac gcgggcaat ggagcagttt ctgcc 45

 <210> 127
 <211> 30
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> primer

 <400> 127
 gccttcctgg ataaaaaatt caagcgatgc 30

 <210> 128
 <211> 31
 <212> DNA
 <213> Artificial Sequence

 <220>
 <223> primer

 <400> 128
 gcacgccttg aattttttat ccaggaaggc g 31

<210> 129
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> synthesized

<400> 129

Pro Lys Lys Lys Arg Lys Val
1 5

<210> 130
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
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<220>
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<222> (4)..(14)
<223> Xaa equals a sequence of any 11 amino acids

<400> 130

Arg Arg Arg Xaa Lys Arg Arg Lys
1 5

<210> 131
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> synthesized

<220>
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<222> (3)..(17)
<223> Xaa equals a sequence of any 15 amino acids

<400> 131

Lys Lys Xaa Lys Lys Arg Lys
1 5

<210> 132
<211> 6
<212> PRT
<213> Artificial Sequence

<220>
<223> synthesized

<400> 132

Lys Arg Lys Arg Arg Pro
1 5

<210> 133

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> synthesized

<400> 133

Pro Lys Lys Asn Arg Leu Arg Arg Lys
1 5

<210> 134

<211> 10

<212> PRT

<213> Artificial Sequence

<220>

<223> synthesized

<220>

<221> MISC_FEATURE

<222> (5)..(24)

<223> Xaa equals a sequence of any 20 amino acids

<400> 134

Lys Arg Gln Arg Xaa Lys Lys Ser Lys Lys
1 5 10

<210> 135

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> synthesized

<400> 135

Pro Ala Ala Lys Arg Val Lys Leu Asp
1 5

<210> 136

<211> 6

<212> PRT

<213> Artificial Sequence

<220>

<223> synthesized

<400> 136

Gln Arg Lys Arg Gln Lys
1 5

<210> 137

<211> 17

<212> PRT

<213> Artificial Sequence

<220>

<223> synthesized

<400> 137

His Arg Ile Glu Glu Lys Arg Lys Arg Thr Tyr Glu Thr Phe Lys Ser
1 5 10 15

Ile

<210> 138

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> synthesized

<400> 138

Lys Lys Lys Tyr Lys Leu Lys
1 5

<210> 139

<211> 7

<212> PRT

<213> Artificial Sequence

<220>

<223> synthesized

<400> 139

Lys Ser Lys Lys Lys Ala Gln
1 5

<210> 140

<211> 9

<212> PRT

<213> Artificial Sequence

<220>

<223> synthesized

<400> 140

Lys Lys Lys Lys Arg Lys Arg Glu Lys
1 5

<210> 141
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> synthesized

<400> 141

Leu Lys Arg Pro Arg Ser Pro Ser Ser
1 5

<210> 142
<211> 13
<212> PRT
<213> Artificial Sequence

<220>
<223> synthesized

<220>
<221> MISC_FEATURE
<222> (4)..(25)
<223> Xaa equals a sequence of any 22 amino acids

<400> 142

Lys Arg Lys Xaa Lys Glu Leu Gln Lys Gln Ile Thr Lys
1 5 10

<210> 143
<211> 9
<212> PRT
<213> Artificial Sequence

<220>
<223> synthesized

<400> 143

Gly Lys Lys Lys Tyr Lys Leu Lys His
1 5

<210> 144
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> synthesized

<400> 144

Lys Lys Lys Tyr Lys Leu Lys
1 5

<210> 145
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> synthesized

<400> 145

Lys Ser Lys Lys Lys Ala Gln
1 5

<210> 146
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
<223> synthesized

<220>
<221> MISC_FEATURE
<222> (4)..(353)
<223> Xaa equals a sequence of any 350 amino acids

<400> 146

Glu Glu Asp Xaa Lys Lys Lys Arg Glu Arg Leu Asp
1 5 10

<210> 147
<211> 25
<212> PRT
<213> Artificial Sequence

<220>
<223> synthesized

<400> 147

Cys Tyr Phe Gln Lys Lys Ala Ala Asn Met Leu Gln Gln Ser Gly Ser
1 5 10 15

Lys Asn Thr Gly Ala Lys Lys Arg Lys
20 25

<210> 148
<211> 12
<212> PRT
<213> Artificial Sequence

<220>
 <223> synthesized

 <220>
 <221> MISC_FEATURE
 <222> (6)..(328)
 <223> Xaa equals a sequence of any 323 amino acids

 <400> 148

Asp Ile Leu Arg Arg Xaa Pro Lys Gln Lys Arg Lys
 1 5 10

<210> 149
 <211> 22
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> synthesized

<400> 149

Ser Ser Asp Asp Glu Ala Thr Ala Asp Ser Gln His Ser Thr Pro Pro
 1 5 10 15

Lys Lys Lys Arg Lys Val
 20

<210> 150
 <211> 12
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> synthesized

<220>
 <221> MISC_FEATURE
 <222> (6)..(14)
 <223> Xaa equals a sequence of any 9 amino acids

<400> 150

Arg Lys Lys Arg Lys Xaa Lys Ala Lys Lys Ser Lys
 1 5 10

<210> 151
 <211> 7
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> synthesized

<220>
 <221> MISC_FEATURE

<222> (3)..(13)
<223> Xaa equals a sequence of any 11 amino acids

<400> 151
Lys Arg Xaa Lys Lys Leu Arg
1 5

<210> 152
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> synthesized

<220>
<221> MISC_FEATURE
<222> (5)..(27)
<223> Xaa equals a sequence of any 22 amino acids

<220>
<221> MISC_FEATURE
<222> (5)..(26)
<223> Xaa equals any amino acid

<400> 152
Arg Arg Pro Ser Xaa Arg Arg Lys Arg Gln Lys
1 5 10

<210> 153
<211> 8
<212> PRT
<213> Artificial Sequence

<220>
<223> synthesized

<220>
<221> MISC_FEATURE
<222> (4)..(14)
<223> Xaa equals a sequence of any 11 amino acids

<400> 153
Arg Arg Arg Xaa Lys Arg Arg Lys
1 5

<210> 154
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> synthesized

<220>

<221> MISC_FEATURE
 <222> (3)..(12)
 <223> Xaa equals a sequence of any 10 amino acids

<400> 154

Lys Arg Xaa Lys Lys Lys Leu
 1 5

<210> 155
 <211> 12
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> synthesized

<220>
 <221> MISC_FEATURE
 <222> (5)..(11)
 <223> Xaa equals a sequence of any 7 amino acids

<400> 155

Arg Lys Arg Lys Xaa Arg Arg Ser Arg Tyr Arg Lys
 1 5 10

<210> 156
 <211> 9
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> synthesized

<400> 156

Met Ile Ser Glu Ala Leu Arg Lys Ala
 1 5

<210> 157
 <211> 5
 <212> PRT
 <213> Artificial Sequence

<220>
 <223> synthesized

<400> 157

Lys Lys Phe Lys Arg
 1 5

<210> 158
 <211> 9
 <212> PRT
 <213> Artificial Sequence

<220>

<223> synthesized

<400> 158

Ala Phe Ser Ala Lys Lys Phe Lys Arg
1 5

<210> 159

<211> 446

<212> PRT

<213> Homo sapiens

<400> 159

Met Arg Thr Leu Asn Thr Ser Ala Met Asp Gly Thr Gly Leu Val Val
1 5 10 15

Glu Arg Asp Phe Ser Val Arg Ile Leu Thr Ala Cys Phe Leu Ser Leu
20 25 30

Leu Ile Leu Ser Thr Leu Leu Gly Asn Thr Leu Val Cys Ala Ala Val
35 40 45

Ile Arg Phe Arg His Leu Arg Ser Lys Val Thr Asn Phe Phe Val Ile
50 55 60

Ser Leu Ala Val Ser Asp Leu Leu Val Ala Val Leu Val Met Pro Trp
65 70 75 80

Lys Ala Val Ala Glu Ile Ala Gly Phe Trp Pro Phe Gly Ser Phe Cys
85 90 95

Asn Ile Trp Val Ala Phe Asp Ile Met Cys Ser Thr Ala Ser Ile Leu
100 105 110

Asn Leu Cys Val Ile Ser Val Asp Arg Tyr Trp Ala Ile Ser Ser Pro
115 120 125

Phe Arg Tyr Glu Arg Lys Met Thr Pro Lys Ala Ala Phe Ile Leu Ile
130 135 140

Ser Val Ala Trp Thr Leu Ser Val Leu Ile Ser Phe Ile Pro Val Gln
145 150 155 160

Leu Ser Trp His Lys Ala Lys Pro Thr Ser Pro Ser Asp Gly Asn Ala

	165		170		175
Thr Ser Leu Ala Glu Thr Ile Asp Asn Cys Asp Ser Ser Leu Ser Arg	180		185		190
Thr Tyr Ala Ile Ser Ser Ser Val Ile Ser Phe Tyr Ile Pro Val Ala	195		200		205
Ile Met Ile Val Thr Tyr Thr Arg Ile Tyr Arg Ile Ala Gln Lys Gln	210		215		220
Ile Arg Arg Ile Ala Ala Leu Glu Arg Ala Ala Val His Ala Lys Asn	225		230		235 240
Cys Gln Thr Thr Thr Gly Asn Gly Lys Pro Val Glu Cys Ser Gln Pro		245		250	255
Glu Ser Ser Phe Lys Met Ser Phe Lys Arg Glu Thr Lys Val Leu Lys		260		265	270
Thr Leu Ser Val Ile Met Gly Val Phe Val Cys Cys Trp Leu Pro Phe		275		280	285
Phe Ile Leu Asn Cys Ile Leu Pro Phe Cys Gly Ser Gly Glu Thr Gln		290		295	300
Pro Phe Cys Ile Asp Ser Asn Thr Phe Asp Val Phe Val Trp Phe Gly	305		310		315 320
Trp Ala Asn Ser Ser Leu Asn Pro Ile Ile Tyr Ala Phe Asn Ala Lys		325		330	335
Lys Phe Lys Arg Phe Ser Thr Leu Leu Gly Cys Tyr Arg Leu Cys Pro		340		345	350
Ala Thr Asn Asn Ala Ile Glu Thr Val Ser Ile Asn Asn Asn Gly Ala		355		360	365
Ala Met Phe Ser Ser His His Glu Pro Arg Gly Ser Ile Ser Lys Glu		370		375	380
Cys Asn Leu Val Tyr Leu Ile Pro His Ala Val Gly Ser Ser Glu Asp	385		390		395 400

Leu Lys Lys Glu Glu Ala Ala Gly Ile Ala Arg Pro Leu Glu Lys Leu
405 410 415

Ser Pro Ala Leu Ser Val Ile Leu Asp Tyr Asp Thr Asp Val Ser Leu
420 425 430

Glu Lys Ile Gln Pro Ile Thr Gln Asn Gly Gln His Pro Thr
435 440 445